

TEEBB" 2004060

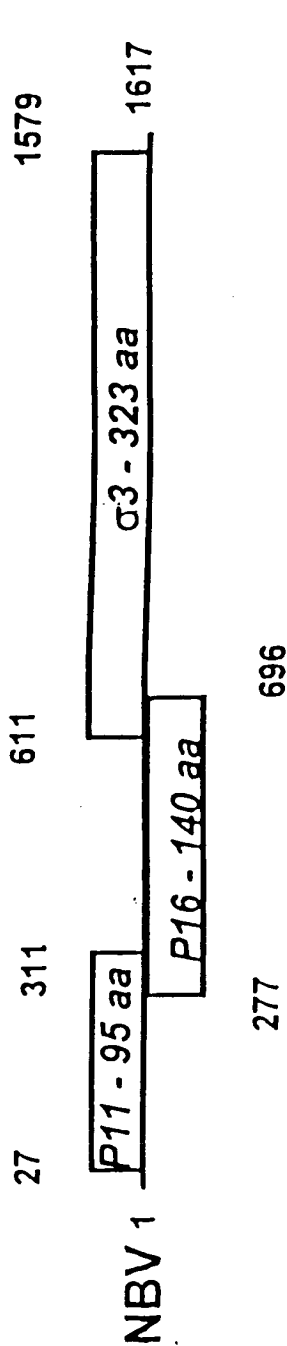


Figure 1A

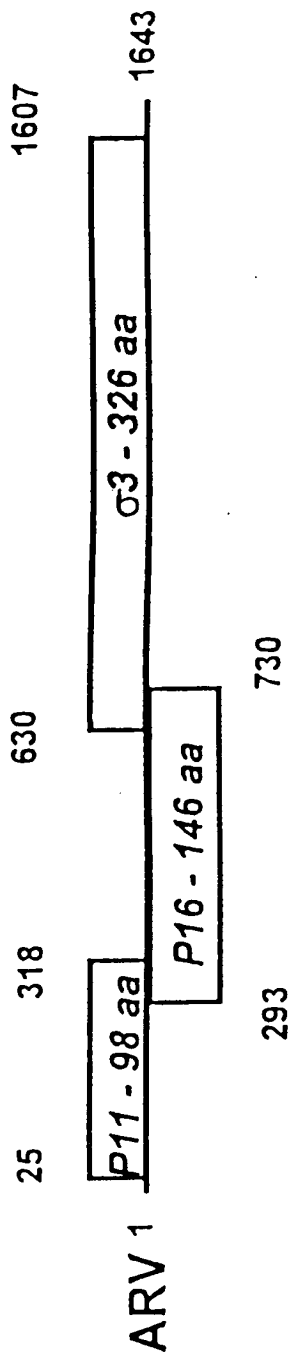


Figure 1B

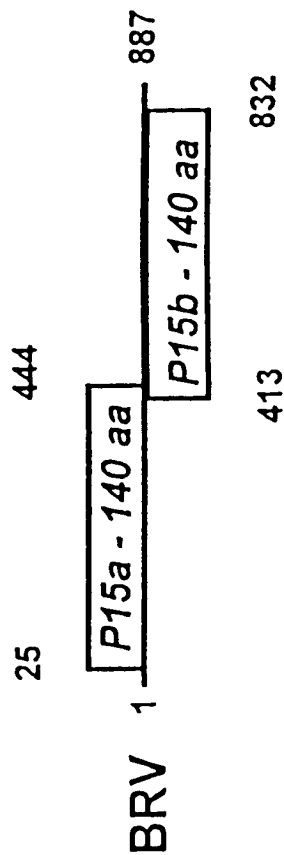


Figure 1C

		*	*	transmembrane
ARV-138	1	MLRMP	PGSCN	GATAIFGNVH COAQNTAGG DLQATSSIIA YWPYLAAGGG <u>FLIIIIIFAIF</u> 61
ARV-176	1	MLRMP	PGSCN	GATAVFGNVH COAQNTAGG DLQATSSIIA YWPYLAAGGG <u>FLIIVIIIFALL</u> 61
NBV	1	....MSSDCA	KIVSVFGSVH	COSSKNSAGG DLQATSVFTT YWPHFAIGGG IIVVILLGLF 57
consensus		-----C-	-----FG-VH	CQ---N-AGG DLQATS---- YWP--A-GGG -----
		**	+	+
ARV-138	62	YCCKAKVKAD	AARSVFHREL	VALSSGKHNA MAPPYDV. 98
ARV-176	62	YCCKAKVKAD	AARSVFHREL	VALSSGKHNA MAPPYDV. 98
NBV	58	YCCYLKWKTS	QVKHTYRREL	IALTRSHVHS TPSGISYV 95
consensus		YCC--K-K--	-----REL	-AL-----

Figure 2

